AUG 2 3 2004 P. TRADEWIND (110)

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847

185

180

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ggacgagttg gagcaactgc tcaagggtgt tgctatgatg aaagagctga ctcttaggac 180
acgagattac cttgtttcct ttggtgaatg catgtctaca agaatatttt ctgcatattt 240
gaataaacta gggaagaagg cacgacagta tgatgctttt gatcttggnt ttataaccac 300
tggacgattt ccacaaatgc cgatatccnc gaacaactta tcctgctgtt gcaaagagct 360
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gaacttgtgc ggcanaactt aggaagggc ggaatgactt gacggcacaa ccatgggaaa 480
cctggggtta agaaaatcag gttggaagat gtaacggttt tgactgtgat caatattatc 540
aaaccggaca ntaccactta ctttgtaggg accgaacttc tnntttggaa agtttgacca 600
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<210> 14
<211> 146
<212> PRT
<213> Triticum aestivum
<220>
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<222> (110)
<223> Xaa=any amino acid
<220>
<221> UNSURE
<222> (131)
<223> Xaa=any amino acid
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<221> UNSURE
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<223> Xaa=any amino acid
Ala Val Ser Cys Gly Ala Pro Lys Ala Ser Glu Ile Tyr Glu Leu Ala
Val Ile Lys Glu Leu His Leu Arg Thr Ile Asp Glu Leu Gly Leu Asp
Ser Ser Ile Val Ser Gly Phe Leu Asp Glu Leu Glu Gln Leu Leu Lys
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Gly Val Ala Met Met Lys Glu Leu Thr Leu Arg Thr Arg Asp Tyr Leu
Val Ser Phe Gly Glu Cys Met Ser Thr Arg Ile Phe Ser Ala Tyr Leu
Asn Lys Leu Gly Lys Lys Ala Arg Gln Tyr Asp Ala Phe Asp Leu Gly
Phe Ile Thr Thr Gly Arg Phe Pro Gln Met Pro Ile Ser Xaa Asn Asn
Leu Ser Cys Cys Cys Lys Glu Leu His Gly Asn Trp Leu Met Thr Leu
        115
                            120
Leu Ser Xaa Tyr Asp Gly Ser Leu Gly Lys Gly Trp Asn Leu Cys Gly
    130
                        135
Xaa Thr
145
<210> 15
<211> 1658
<212> DNA
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                                                                   120
tcaqqttttt tggacgagtt ggagcaactg ctcaagggtg ttgctatgat gaaagagctg
                                                                   180
                                                                   240
actettagga cacqaqatta cettqtttcc tttggtgaat gcatgtctac aagaatattt
                                                                   300
tctqcatatt tqaataaact agggaaqaag gcacgacagt atgatgcttt tgatcttggc
tttataacca ctgacgattt cacaaatgcc gatattctcg aagcaactta tcctgctgtt
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gcaaagaggc tacatggaga ttggattgat gacctgcta ttcctatagt gactggtttc
                                                                  420
cttgggaagg gatggaaatc ttgtgcggtc acaacgttag gaaggggcgg cagtgacttg
                                                                  480
accqctacaa ccattqqcaa aqccttqqqq ttaaqaqaaa ttcaqqtttq qaaqqatqta
                                                                   540
gacggtgtgt tgacgtgtga tccaaatatt tatgcaaacg cggtaccagt accctacttg
                                                                   600
acttttgatg aggcagctga acttgcttat tttggtgcac aggttttgca tccccaatcc
                                                                   660
                                                                   720
atgcgaccag ccagggaagg tggtatccca gttcgagtga agaactcata taaccgtcat
gcacctggca ctgtgatcac taaaacaaga gatatgcgca agagcatatt aaccagcatt
                                                                   780
                                                                   840
gtcctgaaat caaatattac catgctggat atagtgagca caaggatgct cggacagtat
ggctttctag caaaggtctt ctcaatattt gaagatttgg gtatctctgt tgattctgtg
                                                                   900
gctactagtg aagtcagcat atcattgaca ctagatccat caaaactgtg gagtcgtgaa
                                                                   960
ttgatccagc aggagcttga tcatgtagtt gaagagcttg aaaagattgc ggttgttcat 1020
ctcctacage acagateaat cattteectg atagggaatg tgeagagate gtetetgatt 1080
cttgagaagg cgttcaatgt tctacgcaga aatggtgtta atgttcagat gatttcgcaa 1140
ggggcgtcca aggtgaacat ctccttggtg gtgaatgaca gcgaggcgaa gcagtgcgtg 1200
caagccctcc actoggcatt ctttgagaac ggtttcttgt cagaagtaga ggaagcggac 1260
cttgcgcaga agagggctcc agtcctagta agctcgaatg gtgccatcaa cggaaactag 1320
tegaegtege tittitetae tieeageaae ggatgegeeg tiettaggit aagagggtga 1380
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tgtatgtgat tgtgaggcgt cctacttgct gaacttaacc attgtgagga gcccctatga 1560
acttateett gggtgtette taccaaatae taaatagtat gtgtgttgtt cetecaaaaa 1620
1658
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<213> Triticum aestivum

<400> 16

Phe Gly Thr Arg Ala Val Ser Cys Gly Ala Pro Lys Ala Ser Glu Ile 1 5 10 15

Tyr Glu Leu Ala Val Ile Lys Glu Leu His Leu Arg Thr Ile Asp Glu
20 25 30

Leu Gly Leu Asp Ser Ser Ile Val Ser Gly Phe Leu Asp Glu Leu Glu 35 40 45

Gln Leu Leu Lys Gly Val Ala Met Met Lys Glu Leu Thr Leu Arg Thr 50 55 60

Arg Asp Tyr Leu Val Ser Phe Gly Glu Cys Met Ser Thr Arg Ile Phe 65 70 75 80

Ser Ala Tyr Leu Asn Lys Leu Gly Lys Lys Ala Arg Gln Tyr Asp Ala 85 90 95

Phe Asp Leu Gly Phe Ile Thr Thr Asp Asp Phe Thr Asn Ala Asp Ile
100 105 110

Leu Glu Ala Thr Tyr Pro Ala Val Ala Lys Arg Leu His Gly Asp Trp
115 120 125

Ile Asp Asp Pro Ala Ile Pro Ile Val Thr Gly Phe Leu Gly Lys Gly 130 135 140

Trp Lys Ser Cys Ala Val Thr Thr Leu Gly Arg Gly Gly Ser Asp Leu 145 150 155 160

Thr Ala Thr Thr Ile Gly Lys Ala Leu Gly Leu Arg Glu Ile Gln Val 165 170 175

Trp Lys Asp Val Asp Gly Val Leu Thr Cys Asp Pro Asn Ile Tyr Ala 180 185 190

Asn Ala Val Pro Val Pro Tyr Leu Thr Phe Asp Glu Ala Ala Glu Leu 195 200 205

Ala Tyr Phe Gly Ala Gln Val Leu His Pro Gln Ser Met Arg Pro Ala 210 215 220

Arg Glu Gly Gly Ile Pro Val Arg Val Lys Asn Ser Tyr Asn Arg His 225 230 235 240

Ala Pro Gly Thr Val Ile Thr Lys Thr Arg Asp Met Arg Lys Ser Ile 245 250 255

Leu Thr Ser Ile Val Leu Lys Ser Asn Ile Thr Met Leu Asp Ile Val 260 265 270

Ser Thr Arg Met Leu Gly Gln Tyr Gly Phe Leu Ala Lys Val Phe Ser 275 280 285

Ile Phe Glu Asp Leu Gly Ile Ser Val Asp Ser Val Ala Thr Ser Glu 290 295 300

Val Ser Ile Ser Leu Thr Leu Asp Pro Ser Lys Leu Trp Ser Arg Glu 305 310 315 320

Leu Ile Gln Gln Glu Leu Asp His Val Val Glu Glu Leu Glu Lys Ile 325 330 335

Ala Val Val His Leu Leu Gln His Arg Ser Ile Ile Ser Leu Ile Gly 340 345 350

Asn Val Gln Arg Ser Ser Leu Ile Leu Glu Lys Ala Phe Asn Val Leu 355 360 365

Arg Arg Asn Gly Val Asn Val Gln Met Ile Ser Gln Gly Ala Ser Lys 370 375 380

Val Asn Ile Ser Leu Val Val Asn Asp Ser Glu Ala Lys Gln Cys Val 385 390 395 400

Gln Ala Leu His Ser Ala Phe Phe Glu Asn Gly Phe Leu Ser Glu Val 405 410 415

Glu Glu Ala Asp Leu Ala Gln Lys Arg Ala Pro Val Leu Val Ser Ser 420 425 430

Asn Gly Ala Ile Asn Gly Asn 435

<210> 17

<211> 564

<212> PRT

<213> Glycine max

<400> 17

Met Ala Ser Ala Leu Gln Gln Leu Gln Gly Val Gln Gly Lys Leu Ala 1 5 10 15

Val Ser Met Ser Val Arg Arg Ser Leu His His Cys Lys Ser Gln Ile 20 25 30

Gly Phe Ala Ala Leu Gly Ala Pro Val Cys Ala Arg Arg Val Trp Gly
35 40 45

Asn Arg Val Ala Phe Ser Val Thr Thr Cys Lys Ala Ser Thr Ser Asp 50 55 60

Val Ile Glu Lys Asn Ala Thr Glu Asn Gly Met Val Ser Ser Glu Gly 65 70 75 80

Glu Thr Ser Phe Thr Cys Val Met Lys Phe Gly Gly Ser Ser Val Ala 85 90 95

Ser Ala Asp Arg Met Lys Glu Val Ala Thr Leu Ile Leu Ser Phe Pro 100 105 110

Glu Glu Arg Pro Ile Val Val Leu Ser Ala Met Gly Lys Thr Thr Asn 115 120 125

Lys Leu Leu Leu Ala Gly Glu Lys Ala Val Ser Cys Gly Val Ile Asn 130 135 140

Val Ser Ser Ile Glu Glu Leu Cys Phe Ile Lys Asp Leu His Leu Arg 150 Thr Val Asp Gln Leu Gly Val Asp Gly Ser Val Ile Ser Lys His Leu 170 Glu Glu Leu Glu Gln Leu Leu Lys Gly Ile Ala Met Met Lys Glu Leu Thr Lys Arg Thr Gln Asp Tyr Leu Val Ser Phe Gly Glu Cys Met Ser Thr Arg Ile Phe Ala Ala Tyr Leu Asn Lys Ile Gly Val Lys Ala Arg Gln Tyr Asp Ala Phe Glu Ile Gly Phe Ile Thr Thr Asp Asp Phe Thr Asn Ala Asp Ile Leu Glu Ala Thr Tyr Pro Ala Val Ala Lys Arg Leu His Gly Asp Trp Leu Ser Asp Pro Ala Ile Ala Ile Val Thr Gly Phe Leu Gly Lys Ala Arg Lys Ser Cys Ala Val Thr Thr Leu Gly Arg Gly Gly Ser Asp Leu Thr Ala Thr Thr Ile Gly Lys Ala Leu Gly Leu Pro Glu Ile Gln Val Trp Lys Asp Val Asp Gly Val Leu Thr Cys Asp Pro Asn Ile Tyr Pro Lys Ala Glu Pro Val Pro Tyr Leu Thr Phe Asp Glu 325 Ala Ala Glu Leu Ala Tyr Phe Gly Ala Gln Val Leu His Pro Gln Ser 345 Met Arg Pro Ala Arg Glu Ser Asp Ile Pro Val Arg Val Lys Asn Ser 360 Tyr Asn Pro Lys Ala Pro Gly Thr Leu Ile Thr Lys Ala Arg Asp Met 375 Ser Lys Ala Val Leu Thr Ser Ile Val Leu Lys Arg Asn Val Thr Met 390 Leu Asp Ile Ala Ser Thr Arg Met Leu Gly Gln Tyr Gly Phe Leu Ala 405 410 Lys Val Phe Ser Ile Phe Glu Glu Leu Gly Ile Ser Val Asp Val Val 420 Ala Thr Ser Glu Val Ser Val Ser Leu Thr Leu Asp Pro Ser Lys Leu Trp Ser Arg Glu Leu Ile Gln Gln Ala Ser Glu Leu Asp His Val Val 450

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Glu Glu Leu Glu Lys Ile Ala Val Val Asn Leu Leu Gln Asn Arg Ser
465
                    470
                                         475
Ile Ile Ser Leu Ile Gly Asn Val Gln Arg Ser Ser Leu Ile Leu Glu
Arg Leu Ser Arg Val Leu Arg Thr Leu Gly Val Thr Val Gln Met Ile
Ser Gln Gly Ala Ser Lys Val Asn Ile Ser Leu Val Val Asn Asp Ser
                             520
Glu Ala Glu Gln Cys Val Arg Ala Leu His Ser Ala Phe Phe Glu Ser
                        535
Glu Leu Ser Glu Leu Glu Met Asp Tyr Lys Asn Gly Asn Gly Ser Val
                    550
Asp Glu Leu Ser
<210> 18
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<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
<400> 18
ctctctgcca tggggaa
                                                                    17
<210> 19
<211> 27
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic oligonucleotide
<400> 19
gactggtacc tcagcccacg agtaggt
                                                                    27
<210> 20
<211> 23
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
<400> 20
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<210> 21
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<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic oligonucleotide
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<210> 22
<211> 32
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<213> Artificial Sequence
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<210> 23
<211> 33
<212> DNA
<213> Artificial Sequence
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<400> 23
ctttgatgga tcaagtaaca cagaaacact aac
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<210> 24
<211> 23
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
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                                                                    23
<210> 25
<211> 11
<212> PRT
<213> Zea maize
<220>
<223> DOMAIN
<400> 25
Thr Ser Glu Val Ser Val Ser Val Ser Leu Asp
<210> 26
<211> 11
<212> PRT
<213> Escherichia coli
<220>
<223> DOMAIN
<400> 26
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Thr Ser Glu Val Ser Val Ala Leu Thr Leu Asp
<210> 27
<211> 12
<212> PRT
<213> Zea maize
<220>
<223> DOMAIN
<400> 27
Ser Ser Arg Met Leu Gly Gln Tyr Gly Phe Leu Ala
<210> 28
<211> 12
<212> PRT
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<220>
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<400> 28
Ser Leu Asn Met Leu His Ser Arg Gly Phe Leu Ala
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